SEQUENCE LISTING

```
<110> Reed, Jennifer Lynne
<120> RECOMBINANT IL-9 ANTIBODIES AND USES
 THEREOF
<130> 10271-112-999
<140> 10/823,253
<141> 2004-04-12
<150> 60/477,797
<151> 2003-06-10
<150> 60/462,259
<151> 2003-04-11
<160> 65
<170> FastSEQ for Windows Version 4.0
<210> 1
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR1
<400> 1
Gly Tyr Thr Phe Thr Gly Tyr Trp Ile Glu
                5
<210> 2
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR2
Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Pro Asn Glu Lys Phe Lys
1
Gly
<210> 3
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR3
<400> 3
Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr
```

```
<210> 4
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR1
<400> 4
Lys Ala Ser Gln His Val Gly Thr His Val Thr
                5
<210> 5
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR2
<400> 5
Ser Thr Ser Tyr Arg Tyr Ser
<210> 6
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR3
<400> 6
Gln His Phe Tyr Ser Tyr Pro Leu Thr
<210> 7
<211> 122
<212> PRT
<213> Artificial Sequence
<223> VH domain of antibody 4D4
<400> 7
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
                                     10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Gly Tyr
                                 25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                             40
Gly Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe
                         55
                                             60
Lys Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
                                         75
                    70
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85
                                     90
```

```
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
                         105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
       115
<210> 8
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 4D4 and 4D4 H2-1 D11
<400> 8
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                                    10
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Gly Thr His
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
Tyr Ser Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
                                        75
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln His Phe Tyr Ser Tyr Pro Leu
               85
                                    90
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
           100
<210> 9
<211> 122
<212> PRT
<213> Artificial Sequence
<223> VH domain of antibody 4D4 H2-1 D11
<400> 9
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
                                    10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Gly Tyr
            20
                                25
                                                    30
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                           4.0
Gly Glu Trp Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe
                        55
                                           60
Lys Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
                    70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85
                                    90
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
                               105
           100
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
        115
                            120
<210> 10
<211> 17
<212> PRT
```

```
<213> Artificial Sequence
<220>
<223> VH CDR2
<400> 10
Glu Trp Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe Lys
                                   10
Gly
<210> 11
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR1
Gly Tyr Thr Phe Thr Tyr Tyr Trp Ile Glu
                5
<210> 12
<211> 13
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR3
<400> 12
Ala Asp Tyr Tyr Gly Ser Asp His Val Lys Phe Asp Tyr
<210> 13
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR1
Leu Ala Ser Gln His Val Gly Thr His Val Thr
<210> 14
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR2
<400> 14
Gly Thr Ser Tyr Arg Tyr Ser
```

```
<210> 15
<211> 122
<212> PRT
<213> Artificial Sequence
<220>
<223> VH domain of antibody 4D4com-XF-9
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
                                    10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Tyr Tyr
                                25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                            40
Gly Glu Trp Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe
                        55
Lys Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                                    90
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp His Val Lys Phe Asp Tyr Trp
           100
                                105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
<210> 16
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 4D4com-XF-9
<400> 16
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                 5
                                    10
Asp Arg Val Thr Ile Thr Cys Leu Ala Ser Gln His Val Gly Thr His
            20
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Ile
        35
                            40
Tyr Gly Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
                                             60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
                                         75
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln His Phe Tyr Asp Tyr Pro Leu
                85
                                    90
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100
<210> 17
<211> 122
<212> PRT
<213> Artificial Sequence
<220>
```

<223> VH domain of antibody 4D4com-2F9

```
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr Gly Tyr
                                25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                           40
Gly Glu Trp Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe
                        55
                                           60
Lys Gly Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr
                   70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                                   90
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp His Val Lys Phe Asp Tyr Trp
                                105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
<210> 18
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 4D4com-2F9
<400> 18
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                5
                                    1.0
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Gly Thr His
            20
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
                                                4.5
Tyr Gly Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
                                            60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
                                        75
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln His Phe Tyr Glu Tyr Pro Leu
                85
                                    90
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100
<210> 19
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR1
<400> 19
Gly Gly Thr Phe Ser Gly Tyr Trp Ile Glu
        5
<210> 20
<211> 9
<212> PRT
<213> Artificial Sequence
```

```
<220>
<223> VL CDR3
<400> 20
Gln Gln Phe Tyr Glu Tyr Pro Leu Thr
<210> 21
<211> 122
<212> PRT
<213> Artificial Sequence
<220>
<223> VH domain of antibody 7F3 and 7F3 22D3
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Gly Tyr
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                           40
Gly Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe
                        55
                                            60
Lys Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
                    70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85
                                    90
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
           100
                            105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
        115
<210> 22
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 7F3
<400> 22
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                                    10
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Gly Thr His
            20
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
Tyr Ser Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
                                            60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
                                        75
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Tyr Glu Tyr Pro Leu
                85
                                    90
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100
                                105
```

<211> 122

<210> 23

```
<220>
<223> VH domain of antibody 71A10
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
                                    10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Gly Tyr
                                25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                            40
Gly Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Pro Asn Glu Lys Phe
                        55
Lys Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
                    70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                85
                                    90
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
           100
                                105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
        115
<210> 24
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 71A10
<400> 24
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                                    10
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Gly Thr His
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
Tyr Ser Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Tyr Glu Tyr Pro Leu
                85
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
<210> 25
<211> 107
<212> PRT
<213> Artificial Sequence
<220>
<223> VL domain of antibody 7F3 22D3
<400> 25
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                                    10
                 5
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Gly Thr His
```

<212> PRT

<213> Artificial Sequence

```
20
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
Tyr Gly Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                                        75
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Tyr Glu Tyr Pro Leu
               85
                                    90
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100
<210> 26
<211> 10
<212> PRT
<213> Artificial Sequence
<223> VH CDR1
<400> 26
Gly Gly Thr Phe Ser Tyr Tyr Trp Ile Glu
<210> 27
<211> 122
<212> PRT
<213> Artificial Sequence
<223> VH domain of antibody 7F3com-2H2
<400> 27
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
                                    10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Tyr Tyr
            20
                                25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                            40
Gly Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Pro Asn Glu Lys Phe
                        55
                                             60
Lys Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
                    70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                                    90
                85
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
           100
                               105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
        115
<210> 28
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 7F3com-2H2
<400> 28
```

```
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                                    10
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Ile Thr His
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
Tyr Gly Thr Ser Tyr Ser Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Tyr Glu Tyr Pro Leu
               85
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
           100
<210> 29
<211> 122
<212> PRT
<213> Artificial Sequence
<220>
<223> VH domain of antibody 7F3com-3H5
<400> 29
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
                                    10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Gly Tyr
            20
                                25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                            40
                                                45
Gly Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Pro Asn Glu Lys Phe
                        55
                                            60
Lys Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
                    70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                                   90
                85
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
          100
                               105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
        115
<210> 30
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 7F3com-3H5
<400> 30
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                 5
                                    10
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Gly Thr His
            20
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
        35
                            40
Tyr Gly Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
                                            60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
                                        75
```

```
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Tyr Glu Tyr Pro Leu
                                   90
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
<210> 31
<211> 122
<212> PRT
<213> Artificial Sequence
<220>
<223> VH domain of antibody 7F3com-3D4
<400> 31
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
                                    10
Ser Val Lys Val Ser Cys Lys Ala Ser Gly Gly Thr Phe Ser Tyr Tyr
                                25
Trp Ile Glu Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met
                            40
Gly Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Pro Asn Glu Lys Phe
                        55
Lys Gly Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr
                    70
                                        75
Met Glu Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys
                                    90
Ala Arg Ala Asp Tyr Tyr Gly Ser Asp Tyr Val Lys Phe Asp Tyr Trp
            100
                                105
Gly Gln Gly Thr Leu Val Thr Val Ser Ser
<210> 32
<211> 107
<212> PRT
<213> Artificial Sequence
<223> VL domain of antibody 7F3com-3D4
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
                                    10
Asp Arg Val Thr Ile Thr Cys Lys Ala Ser Gln His Val Ile Thr His
                                25
Val Thr Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile
                            40
Tyr Gly Thr Ser Tyr Arg Tyr Ser Gly Val Pro Ser Arg Phe Ser Gly
                        55
                                            60
Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Pro
                    70
                                         75
Glu Asp Phe Ala Thr Tyr Tyr Cys Gln Gln Phe Tyr Glu Tyr Pro Leu
                                    90
                85
Thr Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
            100
<210> 33
<211> 25
<212> PRT
<213> Artificial Sequence
```

```
<220>
<223> VH framework region 1
<400> 33
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ala
Ser Val Lys Val Ser Cys Lys Ala Ser
<210> 34
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 2
<400> 34
Trp Val Arg Gln Ala Pro Gly Gln Gly Leu Glu Trp Met Gly
                 5
<210> 35
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 3
<400> 35
Arg Val Thr Met Thr Arg Asp Thr Ser Thr Ser Thr Val Tyr Met Glu
                 5
                                     10
Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg
<210> 36
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 4
<400> 36
Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ser
<210> 37
<211> 25
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 1
<400> 37
Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Ser
```

```
15
Ser Val Lys Val Ser Cys Lys Ala Ser
<210> 38
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 3
<400> 38
Arg Val Thr Ile Thr Ala Asp Glu Ser Thr Ser Thr Ala Tyr Met Glu
                                    10
Leu Ser Ser Leu Arg Ser Glu Asp Thr Ala Val Tyr Tyr Cys Ala Arg
<210> 39
<211> 23
<212> PRT
<213> Artificial Sequence
<220>
<223> VL framework region 1
<400> 39
Asp Ile Gln Met Thr Gln Ser Pro Ser Ser Leu Ser Ala Ser Val Gly
1
                5
Asp Arg Val Thr Ile Thr Cys
            20
<210> 40
<211> 15
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 2
<400> 40
Trp Tyr Gln Gln Lys Pro Gly Lys Ala Pro Lys Leu Leu Ile Tyr
<210> 41
<211> 32
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 3
<400> 41
Gly Val Pro Ser Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr
                 5
                                     10
Leu Thr Ile Ser Ser Leu Gln Pro Glu Asp Phe Ala Thr Tyr Tyr Cys
            20
                                                     30
                                 25
```

```
<210> 42
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> VH framework region 4
<400> 42
Phe Gly Gly Gly Thr Lys Val Glu Ile Lys
               5
<210> 43
<211> 591
<212> DNA
<213> Artificial Sequence
<220>
<223> VH domain of 7F3com-2H2
ccgctgtcaa gatgcttctg gccatggtcc ttacctctgc cctgctcctg tgctccgtgg 60
caggccaggg gtgtccaacc ttggcgggga tcctggacat caacttcctc atcaacaaga 120
tgcaggaaga tccagcttcc aagtgccact gcagtgctaa tgtgaccagt tgtctctgtt 180
tgggcattcc ctctgacaac tgcaccagac catgcttcag tgagagactg tctcagatga 240
ccaataccac catgcaaaca agatacccac tgattttcag tcgggtgaaa aaatcagttg 300
tgagagggat gagaggcaag atatgaagat gaaatattat ttatcctatt tattaaattt 480
aaaaagcttt ctctttaagt tgctacaatt taaaaatcaa gtaagctact ctaaatcagt 540
atcagttgtg attatttgtt taacattgta tgtctttatt ttgaaataaa t
<210> 44
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> VH CDR1
                                                           30
ggaggcacct tcagctatta ctggatagag
<210> 45
<211> 51
<212> DNA
<213> Artificial Sequence
<220>
<223> VH CDR2
<400> 45
                                                          51
gagattttac ctggaagtgg tactactaac ccgaatgaga agttcaaggg c
<210> 46
<211> 39
<212> DNA
<213> Artificial Sequence
<220>
```

```
<223> VH CDR3
<400> 46
gcggattact acggtagtga ttacgtcaag tttgactac
                                                                    39
<210> 47
<211> 321
<212> DNA
<213> Artificial Sequence
<223> VL domain of 7F3com-2H2
<400> 47
qacatccaga tgacccagtc tccatcctcc ctgtctgcat ctgtaggaga cagagtcacc 60
atcacttgca aggcaagtca gcatgtgatt actcatgtaa cctggtatca gcagaaacca 120
gggaaagccc ctaagctcct gatctatggg acatcctaca gctacagtgg ggtcccatca 180
aggttcagtg gcagtggatc tgggacagat ttcactctca ccatcagcag tctgcaacct 240
gaagattttg caacttatta ctgtcagcaa ttttacgagt atcctctcac gttcggcgga 300
gggaccaagg tggagatcaa a
<210> 48
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> VL CDR1
<400> 48
aaggcaagtc agcatgtgat tactcatgta acc
                                                                    33
<210> 49
<211> 15
<212> DNA
<213> Artificial Sequence
<220>
<223> VL CDR2
<400> 49
                                                                    15
gggacatcct acagc
<210> 50
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> VL CDR3
<400> 50
                                                                    27
cagcaatttt acgagtatcc tctcacg
<210> 51
<211> 591
<212> DNA
<213> Homo sapiens
<400> 51
ccgctgtcaa gatgcttctg gccatggtcc ttacctctgc cctgctcctg tgctccgtgg 60
caggccaggg gtgtccaacc ttggcgggga tcctggacat caacttcctc atcaacaaga 120
```

```
tgcaggaaga tccagcttcc aagtgccact gcagtgctaa tgtgaccagt tgtctctgtt 180
tgggcattcc ctctgacaac tgcaccagac catgcttcag tgagagactg tctcagatga 240
ccaataccac catgcaaaca agatacccac tgattttcag tcgggtgaaa aaatcagttg 300
tgagagggat gagaggcaag atatgaagat gaaatattat ttatcctatt tattaaattt 480
aaaaagettt etetttaagt tgetacaatt taaaaateaa gtaagetaet etaaateagt 540
atcagttgtg attatttgtt taacattgta tgtctttatt ttgaaataaa t
<210> 52
<211> 144
<212> PRT
<213> Homo sapiens
<400> 52
Met Leu Leu Ala Met Val Leu Thr Ser Ala Leu Leu Cys Ser Val
Ala Gly Gln Gly Cys Pro Thr Leu Ala Gly Ile Leu Asp Ile Asn Phe
                              25
Leu Ile Asn Lys Met Gln Glu Asp Pro Ala Ser Lys Cys His Cys Ser
Ala Asn Val Thr Ser Cys Leu Cys Leu Gly Ile Pro Ser Asp Asn Cys
                      55
Thr Arg Pro Cys Phe Ser Glu Arg Leu Ser Gln Met Thr Asn Thr Thr
                  70
                                     75
Met Gln Thr Arg Tyr Pro Leu Ile Phe Ser Arg Val Lys Lys Ser Val
                                 90
Glu Val Leu Lys Asn Asn Lys Cys Pro Tyr Phe Ser Cys Glu Gln Pro
           100
                             105
                                                110
Cys Asn Gln Thr Thr Ala Gly Asn Ala Leu Thr Phe Leu Lys Ser Leu
       115
                          120
                                             125
Leu Glu Ile Phe Gln Lys Glu Lys Met Arg Gly Met Arg Gly Lys Ile
   130
                      135
<210> 53
<211> 808
<212> PRT
<213> Homo sapiens
<400> 53
Met Ala Glu Leu Leu Ala Ser Ala Gly Ser Ala Cys Ser Trp Asp Phe
                                  10
Pro Arg Ala Pro Pro Ser Phe Pro Pro Ala Ala Ser Arg Gly Gly
           20
                              25
Leu Gly Gly Thr Arg Ser Phe Arg Pro His Arg Gly Ala Glu Ser Pro
                          40
                                             4.5
Arg Pro Gly Arg Asp Arg Asp Gly Val Arg Val Pro Met Ala Ser Ser
                      55
                                         60
Arg Cys Pro Ala Pro Arg Gly Cys Arg Cys Leu Pro Gly Ala Ser Leu
                  70
                                     75
Ala Trp Leu Gly Thr Val Leu Leu Leu Ala Asp Trp Val Leu Leu
               8.5
                                  90
Arg Thr Ala Leu Pro Arg Ile Phe Ser Leu Leu Val Pro Thr Ala Leu
                              105
           100
                                                110
Pro Leu Leu Arg Val Trp Ala Val Gly Leu Ser Arg Trp Ala Val Leu
       115
                          120
                                             125
Trp Leu Gly Ala Cys Gly Val Leu Arg Ala Thr Val Gly Ser Lys Ser
                      135
                                         140
Glu Asn Ala Gly Ala Gln Gly Trp Leu Ala Ala Leu Lys Pro Leu Ala
```

155

150

145

```
Ala Ala Leu Gly Leu Ala Leu Pro Gly Leu Ala Leu Phe Arg Glu Leu
                             170
Ile Ser Trp Gly Ala Pro Gly Ser Ala Asp Ser Thr Arg Leu Leu His
           180
                               185
Trp Gly Ser His Pro Thr Ala Phe Val Val Ser Tyr Ala Ala Ala Leu
                           200
Pro Ala Ala Leu Trp His Lys Leu Gly Ser Leu Trp Val Pro Gly
                       215
                                           220
Gly Gln Gly Gly Ser Gly Asn Pro Val Arg Arg Leu Leu Gly Cys Leu
                   230
                                       235
Gly Ser Glu Thr Arg Arg Leu Ser Leu Phe Leu Val Leu Val Val Leu
               245
                                   250
Ser Ser Leu Gly Glu Met Ala Ile Pro Phe Phe Thr Gly Arg Leu Thr
           260
                               265
Asp Trp Ile Leu Gln Asp Gly Ser Ala Asp Thr Phe Thr Arg Asn Leu
        275
                           280
                                                285
Thr Leu Met Ser Ile Leu Thr Ile Ala Ser Ala Val Leu Glu Phe Val
                       295
Gly Asp Gly Ile Tyr Asn Asn Thr Met Gly His Val His Ser His Leu
                   310
                                       315
Gln Gly Glu Val Phe Gly Ala Val Leu Arg Gln Glu Thr Glu Phe Phe
                325
                                   330
Gln Gln Asn Gln Thr Gly Asn Ile Met Ser Arg Val Thr Glu Asp Thr
            340
                               345
Ser Thr Leu Ser Asp Ser Leu Ser Glu Asn Leu Ser Leu Phe Leu Trp
                           360
        355
Tyr Leu Val Arg Gly Leu Cys Leu Leu Gly Ile Met Leu Trp Gly Ser
                        375
                                           380
Val Ser Leu Thr Met Val Thr Leu Ile Thr Leu Pro Leu Leu Phe Leu
                                       395
                   390
Leu Pro Lys Lys Val Gly Lys Trp Tyr Gln Leu Leu Glu Val Gln Val
               405
                                   410
Arg Glu Ser Leu Ala Lys Ser Ser Gln Val Ala Ile Glu Ala Leu Ser
           420
                                425
                                                   430
Ala Met Pro Thr Val Arg Ser Phe Ala Asn Glu Glu Gly Glu Ala Gln
       435
                           440
                                               445
Lys Phe Arg Glu Lys Leu Gln Glu Ile Lys Thr Leu Asn Gln Lys Glu
                       455
                                           460
Ala Val Ala Tyr Ala Val Asn Ser Trp Thr Thr Ser Ile Ser Gly Met
                   470
                                       475
Leu Leu Lys Val Gly Ile Leu Tyr Ile Gly Gly Gln Leu Val Thr Ser
                                   490
                485
                                                        495
Gly Ala Val Ser Ser Gly Asn Leu Val Thr Phe Val Leu Tyr Gln Met
           500
                                505
Gln Phe Thr Gln Ala Val Glu Val Leu Leu Ser Ile Tyr Pro Arg Val
        515
                           520
                                                525
Gln Lys Ala Val Gly Ser Ser Glu Lys Ile Phe Glu Tyr Leu Asp Arg
                        535
                                           540
Thr Pro Arg Cys Pro Pro Ser Gly Leu Leu Thr Pro Leu His Leu Glu
                   550
                                        555
Gly Leu Val Gln Phe Gln Asp Val Ser Phe Ala Tyr Pro Asn Arg Pro
                                    570
                565
Asp Val Leu Val Leu Gln Gly Leu Thr Phe Thr Leu Arg Pro Gly Glu
           580
                                585
Val Thr Ala Leu Val Gly Pro Asn Gly Ser Gly Lys Ser Thr Val Ala
                           600
                                               605
Ala Leu Leu Gln Asn Leu Tyr Gln Pro Thr Gly Gly Gln Leu Leu
                        615
                                            620
Asp Gly Lys Pro Leu Pro Gln Tyr Glu His Arg Tyr Leu His Arg Gln
                   630
                                        635
Val Ala Ala Val Gly Gln Glu Pro Gln Val Phe Gly Arg Ser Leu Gln
                645
                                    650
```

```
Glu Asn Ile Ala Tyr Gly Leu Thr Gln Lys Pro Thr Met Glu Glu Ile
                                665
Thr Ala Ala Ala Val Lys Ser Gly Ala His Ser Phe Ile Ser Gly Leu
                            680
                                                685
Pro Gln Gly Tyr Asp Thr Glu Val Asp Glu Ala Gly Ser Gln Leu Ser
                        695
                                            700
Gly Gly Gln Arg Gln Ala Val Ala Leu Ala Arg Ala Leu Ile Arg Lys
                    710
                                        715
Pro Cys Val Leu Ile Leu Asp Asp Ala Thr Ser Ala Leu Asp Ala Asn
                725
                                    730
Ser Gln Leu Gln Val Glu Gln Leu Leu Tyr Glu Ser Pro Glu Arg Tyr
                                745
Ser Arg Ser Val Leu Leu Ile Thr Gln His Leu Ser Leu Val Glu Gln
                            760
        755
Ala Asp His Ile Leu Phe Leu Glu Gly Gly Ala Ile Arg Glu Gly Gly
                                             780
                        775
Thr His Gln Gln Leu Met Glu Lys Lys Gly Cys Tyr Trp Ala Met Val
                                        795
                    790
Gln Ala Pro Ala Asp Ala Pro Glu
                805
<210> 54
<211> 140
<212> PRT
<213> Homo sapiens
<400> 54
Met Val Leu Thr Ser Ala Leu Leu Cys Ser Val Ala Gly Gln Gly
                 5
                                    10
Cys Pro Thr Leu Ala Gly Ile Leu Asp Ile Asn Phe Leu Ile Asn Lys
                                25
Met Gln Glu Asp Pro Ala Ser Lys Cys His Cys Ser Ala Asn Val Thr
                             40
                                                 45
Ser Cys Leu Cys Leu Gly Ile Pro Ser Asp Asn Cys Thr Arg Pro Cys
                         55
                                             60
Phe Ser Glu Arg Leu Ser Gln Met Thr Asn Thr Thr Met Gln Thr Arg
                    70
                                         75
Tyr Pro Leu Ile Phe Ser Arg Val Lys Lys Ser Val Glu Val Leu Lys
                                     90
Asn Asn Lys Cys Pro Tyr Phe Ser Cys Glu Gln Pro Cys Asn Gln Thr
                                 105
                                                     110
Thr Ala Gly Asn Ala Leu Thr Phe Leu Lys Ser Leu Leu Glu Ile Phe
                             120
Gln Lys Glu Lys Met Arg Gly Met Arg Gly Lys Ile
                         135
<210> 55
<211> 2171
<212> DNA
<213> Homo sapiens
<400> 55
agcagetetg taatgegett gtggttteag atgtgggegg eetgtgtgaa eetgtegtge 60
aaagctcacg tcaccaactg ctgcagttat ctcctgaatc aggctgaggg tctttgctgt 120
gcacccagag atagttgggt gacaaatcac ctccaggttg gggatgcctc agacttgtga 180
tgggactggg cagatgcatc tgggaaggct ggaccttgga gagtgaggcc ctgaggcgag 240
acatgggcac ctggctcctg gcctgcatct gcatctgcac ctgtgtctgc ttgggagtct 300
ctgtcacagg ggaaggacaa gggccaaggt ctagaacctt cacctgcctc accaacaaca 360
ttctcaggat cgattgccac tggtctgccc cagagetggg acagggctcc agcccctggc 420
tectetteae cageaaceag geteetggeg geacacataa gtgeatettg eggggeagtg 480
```

```
agtgcaccgt cgtgctgcca cctgaggcag tgctcgtgcc atctgacaat ttcaccatca 540
ctttccacca ctgcatgtct gggagggagc aggtcagcct ggtggacccg gagtacctgc 600
cccggagaca cgttaagctg gacccgccct ctgacttgca gagcaacatc agttctggcc 660
actgcatcct gacctggagc atcagtcctg ccttggagcc aatgaccaca cttctcagct 720
atgagetgge etteaagaag eaggaagag eetgggagea ggeeeageae agggateaea 780
ttgtcggggt gacctggctt atacttgaag cctttgagct ggaccctggc tttatccatg 840
aggccaggct gcgtgtccag atggccacac tggaggatga tgtggtagag gaggagcgtt 900
atacaggcca gtggagtgag tggagccagc ctgtgtgctt ccaggctccc cagagacaag 960
gccctctgat cccaccctgg gggtggccag gcaacaccct tgttgctgtg tccatctttc 1020
tectgetgae tggeeegaee taceteetgt teaagetgte geeeagggtg aagagaatet 1080
tctaccagaa cgtgccctct ccagcgatgt tcttccagcc cctctacagt gtacacaatg 1140
ggaacttcca gacttggatg ggggcccacg gggccggtgt gctgttgagc caggactgtg 1200
ctggcacccc acagggagcc ttggagccct gcgtccagga ggccactgca ctgctcactt 1260
gtggcccagc gcgtccttgg aaatctgtgg ccctggagga ggaacaggag ggccctggga 1320
ccaggetece ggggaacetg ageteagagg atgtgetgee ageagggtgt aeggagtgga 1380
gggtacagac gcttgcctat ctgccacagg aggactgggc ccccacgtcc ctgactaggc 1440
cggctccccc agactcagag ggcagcagga gcagcagcag cagcagcagc agcaacaaca 1500
acaactactg tgccttgggc tgctatgggg gatggcacct ctcagccctc ccaggaaaca 1560
cacagagete tgggeceate ceagecetgg cetgtggeet ttettgtgae cateagggee 1620
tggagaccca gcaaggagtt gcctgggtgc tggctggtca ctgccagagg cctgggctgc 1680
atgaggacct ccagggcatg ttgctccctt ctgtcctcag caaggctcgg tcctggacat 1740
tctaggtccc tgactcgcca gatgcatcat gtccattttg ggaaaatgga ctgaagtttc 1800
tggagccctt gtctgagact gaacctcctg agaaggggcc cctagcagcg gtcagaggtc 1860
ctgtctggat ggaggctgga ggctccccc tcaacccctc tgctcagtgc ctgtggggag 1920
cagoctetac cetcageate etggecacaa gttetteett ecattgteee ttttettat 1980
ccctgacctc tctgagaagt ggggtgtggt ctctcagctg ttctgccctc atacccttaa 2040
agggccagcc tgggcccagt ggacacaggt aaggcaccat gaccacctgg tgtgacctct 2100
ctgtgcctta ctgaggcacc tttctagaga ttaaaagggg cttgatggct gttaaaaaaa 2160
                                                                  2171
aaaaaaaaa a
```

<210> 56 <211> 2175 <212> DNA <213> Homo sapiens

<400> 56 agcagetetg taatgegett gtggttteag atgtgggegg eetgtgtgaa eetgtegtge 60 aaagctcacg tcaccaactg ctgcagttat ctcctgaatc aggctgaggg tctttgctgt 120 gcacccagag atagttgggt gacaaatcac ctccaggttg gggatgcctc agacttgtga 180 tgggactggg cagatgcatc tgggaagtaa ctgctgcaag aacggacaga cactgctgca 240 gagaacttgc cacggtgttt catgctgtgg ctggtggttc caggctgcac gctccattct 300 aggaaagggg ccctcagccc agtcccttgc aggctggacc ttggagagtg aggccctgag 360 gcgagacatg ggcacctggc tcctggcctg catctgcatc tgcacctgtg tctgcttggg 420 agtetetgte acaggggaag gacaagggee aaggtetaga acetteacet geeteaceaa 480 caacattete aggategatt gecaetggte tgeeceagag etgggacagg getecageee 540 ctggctcctc ttcaccaggc tcctggcggc acacataagt gcatcttgcg gggcagtgag 600 tgcaccgtcg tgctgccacc tgaggcagtg ctcgtgccat ctgacaattt caccatcact 660 ttccaccact gcatgtctgg gagggagcag gtcagcctgg tggacccgga gtacctgccc 720 cggagacacg agcaacatca gttctggcca ctgcatcctg acctggagca tcagtcctgc 780 cttggagcca atgaccacac ttctcagcta tgagctggcc ttcaagaagc aggaagaggc 840 ctgggagcag gcccagcaca gggatcacat tgtcggggtg acctggctta tacttgaagc 900 ctttgagctg gaccctggct ttatccatga ggccaggctg cgtgtccaga tggccacact 960 ggaggatgat gtggtagagg aggagcgtta tacaggccag tggagtgagt ggagccagcc 1020 tgtgtgcttc caggctcccc agagacaagg ccctctgatc ccaccctggg ggtggccagg 1080 caacaccett gttgetgtgt ceatetttet cetgetgaet ggeeegaeet aceteetgtt 1140 caaqctqtcq cccaqacttq qatqqqqqcc cacggggccg gtgtgctgtt gagccaggac 1200 tgtgctggca ccccacaggg agccttggag ccctgcgtcc aggaggccac tgcactgctc 1260 acttgtggcc cagcgcgtcc ttggaaatct gtggccctgg aggaggaaca ggagggccct 1320 gggaccagge teceggggaa eetgagetea gaggatgtge tgeeageagg gtgtaeggag 1380 tggagggtac agacgettge etatetgeca caggaggaet gggeececae gteectgaet 1440 aggccggctc ccccagactc agagggcagc aggagcagca gcagcagcag cagcagcaac 1500

```
aacaacaact actgtgcctt gggctgctat gggggatggc acctctcaqc cctcccagga 1560
aacacacaga gctctgggcc catcccagcc ctggcctgtg gcctttcttg tgaccatcag 1620
ggcctggaga cccagcaagg agttgcctgg gtgctggctg gtcactgcca gaggcctggg 1680
ctgcatgagg acctccaggg catgttgctc ccttctgtcc tcagcaaggc tcggtcctgg 1740
acattetagg tecetgaete geeagatgea teatgteeat ttttgggaaaa tggaetgaag 1800
tttctggagc ccttgtctga gactgaacct cctgagaagg ggcccctagc agcggtcaga 1860
ggtcctgtct ggatggaggc tggaggctcc cccctcaacc cctctgctca gtgcctgtgg 1920
ggagcagcct ctaccctcag catcctggcc acaagttctt ccttccattg tcccttttct 1980
ttatccctga cctctctgag aagtggggtg tggtctctca gctgttctgc cctcataccc 2040
ttaaagggcc agcctgggcc cagtggacac aggtaaggca ccatgaccac ctggtgtgac 2100
ctctctgtgc cttactgagg cacctttcta gagattaaaa ggggcttgat ggctgttaaa 2160
                                                                  2175
aaaaaaaaa aaaaa
<210> 57
<211> 1451
<212> DNA
<213> Homo sapiens
<400> 57
gaagagcaag cgccatgttg aagccatcat taccattcac atccctctta ttcctgcage 60
tgcccctgct gggagtgggg ctgaacacga caattctgac gcccaatggg aatgaagaca 120
ccacagetga tttetteetg accaetatge ccaetgaete ceteagtgtt tecaetetge 180
ccctcccaga ggttcagtgt tttgtgttca atgtcgagta catgaattgc acttggaaca 240
gcagetetga geeceageet accaacetea etetgeatta ttggtacaag aacteggata 300
atgataaagt ccagaagtgc agccactatc tattctctga agaaatcact tctggctgtc 360
agttgcaaaa aaaggagatc cacctctacc aaacatttgt tgttcagctc caggacccac 420
gggaacccag gagacaggcc acacagatgc taaaactgca gaatctggtg atcccctggg 480
ctccagagaa cctaacactt cacaaactga gtgaatccca gctagaactg aactggaaca 540
acagattett gaaccactgt ttggagcact tggtgeagta ceggactgae tgggaccaca 600
gctggactga acaatcagtg gattatagac ataagttctc cttgcctagt gtggatgggc 660
agaaacgcta cacgtttcgt gttcggagcc gctttaaccc actctgtgga agtgctcagc 720
attggagtga atggagccac ccaatccact gggggagcaa tacttcaaaa gagaatcctt 780
teetgtttge attggaagee gtggttatet etgttggete eatgggattg attateagee 840
ttctctgtgt gtatttctgg ctggaacgga cgatgccccg aattcccacc ctgaagaacc 900
tagaggatct tgttactgaa taccacggga acttttcggc ctggagtggt gtgtctaagg 960
gactggctga gagtctgcag ccagactaca gtgaacgact ctgcctcgtc agtgagattc 1020
ccccaaaagg aggggcctt ggggaggggc ctggggcctc cccatgcaac cagcatagcc 1080
cctactgggc cccccatgt tacaccctaa agcctgaaac ctgaacccca atcctctgac 1140
agaagaaccc cagggtcctg tagccctaag tggtactaac tttccttcat tcaacccacc 1200
tgcgtctcat actcacctca ccccactgtg gctgatttgg aattttgtgc ccccatgtaa 1260
gcaccccttc atttggcatt ccccacttga gaattaccct tttgccccga acatgttttt 1320
cttctccctc agtctggccc ttccttttcg caggattctt cctccctccc tctttccctc 1380
ccttcctctt tccatctacc ctccgattgt tcctgaaccg atgagaaata aagtttctgt 1440
                                                                   1451
tgataatcat c
<210> 58
<211> 521
<212> PRT
<213> Homo sapiens
<400> 58
Met Gly Leu Gly Arg Cys Ile Trp Glu Gly Trp Thr Leu Glu Ser Glu
                                    10
Ala Leu Arg Arg Asp Met Gly Thr Trp Leu Leu Ala Cys Ile Cys Ile
            20
                                25
Cys Thr Cys Val Cys Leu Gly Val Ser Val Thr Gly Glu Gly Gln Gly
                                                45
                            40
Pro Arg Ser Arg Thr Phe Thr Cys Leu Thr Asn Asn Ile Leu Arg Ile
                        55
                                             60
Asp Cys His Trp Ser Ala Pro Glu Leu Gly Gln Gly Ser Ser Pro Trp
```

```
70
                                       75
Leu Leu Phe Thr Ser Asn Gln Ala Pro Gly Gly Thr His Lys Cys Ile
                                   90
Leu Arg Gly Ser Glu Cys Thr Val Val Leu Pro Pro Glu Ala Val Leu
           100
                               105
Val Pro Ser Asp Asn Phe Thr Ile Thr Phe His His Cys Met Ser Gly
                           120
Arg Glu Gln Val Ser Leu Val Asp Pro Glu Tyr Leu Pro Arg Arg His
                       135
                                           140
Val Lys Leu Asp Pro Pro Ser Asp Leu Gln Ser Asn Ile Ser Ser Gly
                                       155
                   150
His Cys Ile Leu Thr Trp Ser Ile Ser Pro Ala Leu Glu Pro Met Thr
               165
                                    170
Thr Leu Leu Ser Tyr Glu Leu Ala Phe Lys Lys Gln Glu Glu Ala Trp
                                185
                                                    190
Glu Gln Ala Gln His Arg Asp His Ile Val Gly Val Thr Trp Leu Ile
                            200
                                                205
Leu Glu Ala Phe Glu Leu Asp Pro Gly Phe Ile His Glu Ala Arg Leu
                        215
                                            220
Arg Val Gln Met Ala Thr Leu Glu Asp Asp Val Val Glu Glu Glu Arg
                   230
                                        235
Tyr Thr Gly Gln Trp Ser Glu Trp Ser Gln Pro Val Cys Phe Gln Ala
               245
                                    250
Pro Gln Arg Gln Gly Pro Leu Ile Pro Pro Trp Gly Trp Pro Gly Asn
                                265
Thr Leu Val Ala Val Ser Ile Phe Leu Leu Leu Thr Gly Pro Thr Tyr
                            280
       275
Leu Leu Phe Lys Leu Ser Pro Arg Val Lys Arg Ile Phe Tyr Gln Asn
                        295
Val Pro Ser Pro Ala Met Phe Phe Gln Pro Leu Tyr Ser Val His Asn
                                        315
                    310
Gly Asn Phe Gln Thr Trp Met Gly Ala His Gly Ala Gly Val Leu Leu
                325
                                    330
Ser Gln Asp Cys Ala Gly Thr Pro Gln Gly Ala Leu Glu Pro Cys Val
                                345
            340
Gln Glu Ala Thr Ala Leu Leu Thr Cys Gly Pro Ala Arg Pro Trp Lys
                            360
Ser Val Ala Leu Glu Glu Glu Glu Gly Pro Gly Thr Arg Leu Pro
                        375
Gly Asn Leu Ser Ser Glu Asp Val Leu Pro Ala Gly Cys Thr Glu Trp
                   390
                                        395
Arg Val Gln Thr Leu Ala Tyr Leu Pro Gln Glu Asp Trp Ala Pro Thr
                405
                                    410
Ser Leu Thr Arg Pro Ala Pro Pro Asp Ser Glu Gly Ser Arg Ser Ser
           420
                               425
Ser Ser Ser Ser Ser Asn Asn Asn Tyr Cys Ala Leu Gly Cys
                           440
                                                445
        435
Tyr Gly Gly Trp His Leu Ser Ala Leu Pro Gly Asn Thr Gln Ser Ser
                       455
                                            460
Gly Pro Ile Pro Ala Leu Ala Cys Gly Leu Ser Cys Asp His Gln Gly
                    470
                                        475
Leu Glu Thr Gln Gln Gly Val Ala Trp Val Leu Ala Gly His Cys Gln
                                                       495
                                    490
                485
Arg Pro Gly Leu His Glu Asp Leu Gln Gly Met Leu Leu Pro Ser Val
                                505
            500
Leu Ser Lys Ala Arg Ser Trp Thr Phe
        515
```

<210> 59

<211> 332

<212> PRT

<213> Homo sapiens

```
Met His Leu Gly Ser Asn Cys Cys Lys Asn Gly Gln Thr Leu Leu Gln
                                   10
Arg Thr Cys His Gly Val Ser Cys Cys Gly Trp Trp Phe Gln Ala Ala
Arg Ser Ile Leu Gly Lys Gly Pro Ser Ala Gln Ser Leu Ala Gly Trp
                           40
Thr Leu Glu Ser Glu Ala Leu Arg Arg Asp Met Gly Thr Trp Leu Leu
                       55
Ala Cys Ile Cys Ile Cys Thr Cys Val Cys Leu Gly Val Ser Val Thr
                   70
Gly Glu Gly Gln Gly Pro Arg Ser Arg Thr Phe Thr Cys Leu Thr Asn
                                   90
               85
Asn Ile Leu Arg Ile Asp Cys His Trp Ser Ala Pro Glu Leu Gly Gln
                               105
           100
Gly Ser Ser Pro Trp Leu Leu Phe Thr Arg Leu Leu Ala Ala His Ile
                           120
Ser Ala Ser Cys Gly Ala Val Ser Ala Pro Ser Cys Cys His Leu Arg
                       135
                                           140
Gln Cys Ser Cys His Leu Thr Ile Ser Pro Ser Leu Ser Thr Thr Ala
                                       155
                   150
Cys Leu Gly Gly Ser Arg Ser Ala Trp Trp Thr Arg Ser Thr Cys Pro
                                  170
               165
                                                       175
Gly Asp Thr Ser Asn Ile Ser Ser Gly His Cys Ile Leu Thr Trp Ser
           180
                               185
                                                  190
Ile Ser Pro Ala Leu Glu Pro Met Thr Thr Leu Leu Ser Tyr Glu Leu
       195
                           200
Ala Phe Lys Lys Gln Glu Glu Ala Trp Glu Gln Ala Gln His Arg Asp
                       215
                                           220
His Ile Val Gly Val Thr Trp Leu Ile Leu Glu Ala Phe Glu Leu Asp
                   230
                                       235
Pro Gly Phe Ile His Glu Ala Arg Leu Arg Val Gln Met Ala Thr Leu
                                  250
               245
Glu Asp Asp Val Val Glu Glu Glu Arg Tyr Thr Gly Gln Trp Ser Glu
                               265
           260
Trp Ser Gln Pro Val Cys Phe Gln Ala Pro Gln Arg Gln Gly Pro Leu
                           280
Ile Pro Pro Trp Gly Trp Pro Gly Asn Thr Leu Val Ala Val Ser Ile
                       295
                                           300
Phe Leu Leu Thr Gly Pro Thr Tyr Leu Leu Phe Lys Leu Ser Pro
                310
                                       315
Arg Leu Gly Trp Gly Pro Thr Gly Pro Val Cys
                325
```

```
<210> 60
<211> 369
<212> PRT
<213> Homo sapiens
```

<400> 60

 Met
 Leu
 Lys
 Pro
 Ser
 Leu
 Pro
 Phe
 Thr
 Ser
 Leu
 Phe
 Leu
 Leu
 Leu
 Phe
 Leu
 Leu
 Phe
 Leu
 Phe
 Leu
 Phe
 Leu
 Phe
 Phe</th

```
70
                                       75
Gln Pro Thr Asn Leu Thr Leu His Tyr Trp Tyr Lys Asn Ser Asp Asn
                                   90
Asp Lys Val Gln Lys Cys Ser His Tyr Leu Phe Ser Glu Glu Ile Thr
                               105
Ser Gly Cys Gln Leu Gln Lys Lys Glu Ile His Leu Tyr Gln Thr Phe
                          120
                                           125
Val Val Gln Leu Gln Asp Pro Arg Glu Pro Arg Arg Gln Ala Thr Gln
                      135
                                           140
Met Leu Lys Leu Gln Asn Leu Val Ile Pro Trp Ala Pro Glu Asn Leu
                  150
                                      155
Thr Leu His Lys Leu Ser Glu Ser Gln Leu Glu Leu Asn Trp Asn Asn
               165
                                   170
Arg Phe Leu Asn His Cys Leu Glu His Leu Val Gln Tyr Arg Thr Asp
           180
                               185
Trp Asp His Ser Trp Thr Glu Gln Ser Val Asp Tyr Arg His Lys Phe
       195
                           200
Ser Leu Pro Ser Val Asp Gly Gln Lys Arg Tyr Thr Phe Arg Val Arg
                       215
                                            220
Ser Arg Phe Asn Pro Leu Cys Gly Ser Ala Gln His Trp Ser Glu Trp
                   230
                                       235
Ser His Pro Ile His Trp Gly Ser Asn Thr Ser Lys Glu Asn Pro Phe
               245
                                    250
Leu Phe Ala Leu Glu Ala Val Val Ile Ser Val Gly Ser Met Gly Leu
                                265
Ile Ile Ser Leu Leu Cys Val Tyr Phe Trp Leu Glu Arg Thr Met Pro
       275
                           280
Arg Ile Pro Thr Leu Lys Asn Leu Glu Asp Leu Val Thr Glu Tyr His
                       295
Gly Asn Phe Ser Ala Trp Ser Gly Val Ser Lys Gly Leu Ala Glu Ser
                   310
                                        315
Leu Gln Pro Asp Tyr Ser Glu Arg Leu Cys Leu Val Ser Glu Ile Pro
               325
                                    330
Pro Lys Gly Gly Ala Leu Gly Glu Gly Pro Gly Ala Ser Pro Cys Asn
                                345
Gln His Ser Pro Tyr Trp Ala Pro Pro Cys Tyr Thr Leu Lys Pro Glu
Thr
<210> 61
<211> 17
<212> PRT
<213> Artificial Sequence
<220>
<223> VH CDR2
<400> 61
Glu Ile Leu Pro Gly Ser Gly Thr Thr Asn Tyr Asn Glu Lys Phe Lys
Gly
<210> 62
<211> 11
<212> PRT
<213> Artificial Sequence
```

<220>

```
<223> VL CDR1
<400> 62
Lys Ala Ser Gln His Val Ile Thr His Val Thr
               5
<210> 63
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR3
<400> 63
Gln His Phe Tyr Asp Tyr Pro Leu Thr
       5
<210> 64
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR3
<400> 64
Gln His Phe Tyr Glu Tyr Pro Leu Thr
<210> 65
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> VL CDR2
<400> 65
Gly Thr Ser Tyr Ser Tyr Ser
```